



## COMPUTER SEARCH RESULTS

\*1: P58810. Lambda/chi-conoto...[gi:20454878] BLink, Links

>gi|20454878|sp|P58810|CXL4\_CONMR Lambda/chi-conotoxin MrIB (Chi-MrIB)  
VGVCCGYKLCHPC

\*2: P58808. Lambda-conotoxin ...[gi:20454876] BLink, Links

>gi|20454876|sp|P58808|CXL2\_CONMR Lambda-conotoxin CMrVIB precursor (Chi-conotoxin  
MrIA) (Chi-MrIA) (mr10a)  
MRCLPVLIILLLTASAPGVVVLPKTEDDVPMSVYNGKSLRGILRNGVCCGYKLCHP  
C

\*3: P58807. Lambda-conotoxin ...[gi:20454875] BLink, Links

>gi|20454875|sp|P58807|CXL1\_CONMR Lambda-conotoxin CMrVIA  
VCCGYKLCHPC

4: P58809. Lambda-conotoxin ...[gi:20454877] Links

>gi|20454877|sp|P58809|CXL3\_CONMR Lambda-conotoxin CMrX  
GICCGVSFCYPC

5: AAG60408. conotoxin scaffold...[gi:12619493] BLink, Links

>gi|12619493|gb|AAG60408.1|AF214980\_1 conotoxin scaffold IX precursor [Conus pennaceus]  
MRCLPVFVILLLTASGPSVDAKVHLKTGDGPLSSFRDNAKSTLQRLQDKSTCCGYR  
MCVPCG

6: CAB95439. hypothetical prot...[gi:9366677] BLink, Links

>gi|9366677|emb|CAB95439.1| hypothetical protein [Trypanosoma brucei]  
MKFLVVDGDWPVTPYTLTRSELVSHSVEEADGGSEGGQLIHGHTLVALCQQQLHTSVV  
RVRAAPDLSTLGLTILHESLSDSTGNIRLVTPKSLTRQVFAVDDLLVHGRLIVFTSRWESL  
ASAAIVGVKSDRREEHELTRRTKSPEETKNSGRGNHDTSVRSPSVFSGRNRTALHKFTC  
AETYFQSIPPATVHLDLLNNLIQAGTIRLDPRVEVYLCTVCGFLPRQTFTSCCGAIMC  
GAECTPLISTSSVMP SVMSER YVCIA CKELPFTSHTAH VSRDAEV MKLV REL RV LHPQL  
ASAASATSSHGMAEKSHASLPFLLHSMGIMP AKRDVG GAAAPV LG SVFPNP

\*7: XP\_371536. similar to tripar...[gi:41124489] BLINK, Domains, Links

>gi|41124489|ref|XP\_371536.1| similar to tripartite motif-containing 43 [Homo sapiens]  
MDSDFSHAFQKEELTCVICLNYLVDPVTICCGHSFCRPLCLSWEEAQSPANCPACREPSP  
KMDFKTNILLKNLVTIARKASLWQFLSSEKQICGTHRQTKKMFCMDKSLLCLLCNSQ  
EHGAHKHPIEEAAEEDREKLLKQMRILWKKIQENQRNLYEERRTAFLLRGDVVLRAQ  
MIRNEYRKLHPVLHKEEKQHLERLNKEYQEIFFQQLQRSWVKMDQSKSHLKEMLYQELM  
EMCHKPEVELLQDLGDIVARSESVLLHMPQPVNPELTAGPITGLVYRLNFRVEISFHFE  
VTNHNIRLFEDVRSWMFRRGPLNSDRSDYFAAWGARVFSFGKHYWELDVDNSCDWAL  
GVCNNSWIRKNSTMVNSEDIFLLLCLKVDNHFNLTTSPVFPHYIEKPLGRGVGVFLDFES  
GSVSFLNVTKSSLIWSYPAGSLTFPVRFYTGHR

\*8: NP\_620155. tripartite motif-...[gi:20270353] BLINK, Domains, Links

>gi|20270353|ref|NP\_620155.1| tripartite motif-containing 43 [Homo sapiens]  
MDSDFSHAFQKEELTCVICLNYLVDPVTICCGHSFCRPLCLSWEEAQSPANCPACREPSP  
KMDFKTNILLKNLVTIARKASLWQFLSSEKQICGTHRQTKKMFCMDKSLLCLLCNSQ  
EHGAHKHPIEEAAEHEKLLKQMRILWKKIQENQRNLYEEGRTAFLWRGNVVLRAQ  
MIRNEYRKLHPVLHKEEKQHLERLNKEYQEIFFQQLQRSWVKMDQSKSHLKEMLYQELM  
EMCHKPDVELLQDLGDIVARSESVLLHMPQPVNPELTAGPITGLVYRLNFRVEISFHFE  
VTNHNIRLFEDVRSWMFRRGPLNSDRSDYFAAWGARVFSFGKHYWELDVDNSCDWAL  
GVCNNSWIRKNSTMVNSEDIFLLLCLKVDNHFNLTTSPVFPHYIEKPLGRGVGVFLDFES  
GSVSFLNVTKSSLIWSYPAGSLTFPVRFYTGHR

9: NP\_041168. Zn-binding protei...[gi:9626898] BLINK, Links

>gi|9626898|ref|NP\_041168.1| Zn-binding protein [Ictalurid herpesvirus 1]  
MELSLYRELPRKRRAVAQPRTRQPPPKVHREDTGIVPGLREVESSVRPDSITNFRRLAALK  
FKNPEVMECGGKGGECECKTRPALMLNCDCDTAKVPICLGCKPRAVRDHPCFRNRTCV  
SCGTIGVVAVVKLGCCGATLCDSCLGADWYPKSRTGTCGYCGAQTLGVHYRPMKIIK  
NTRPYIPAETPIPKVQRHISARLLEEIRRCQLSRSEIVSLFGDHGSMMEYPTAVPAHLLAP  
KMIGYSDISKAFADDLPRCSRAMCAANGVPGWRCGFQAPVLFPGMDMCLLCVVRAQ  
EQTITQLVLNGTIWEDAPIAGTYHMSVWLDPTTSVTPVDISAITFAVEFCGNLGSHP  
EKFYRLSDLTCRIRWTAGGFLDPESLMVPEKERLKNTASAHSNNSH